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    Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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hypothetical protein PAB1174 - Pyrococcus abyssi (strain Orsay) C;Species: Pyrococcus abyssi C;Date: 20-Aug-1999 #sequence_revision 20-Aug-1999 #text_change C;Accession: H75029 R;anonymous, Genoscope submitted to the EMBL Data Library, July 1999 A;Description: Pyrococcus abyssi genome sequence: insights into A;Reference number: A75001 A;Accession: H75029
                                                                                                                                                                                                                                             R; Ainscough, R. submitted to the EMBL Data Library, A. Paference number: Z19629
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H75029
                                                                                                                                            A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-351 <WIL>
A;Cross-references: EMBL:Z81554; PIDN:CAB04503.1; GSPDB:GN00023; CESP:F57G4.4
A;Experimental source: clone F57G4
                                                                                                                                                                                                                                                                                                            hypothetical protein F57G4.4 - Caenorhabditis elegans
C; Species: Caenorhabditis elegans
C; Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 15-Oct-1999
C; Accession: T22861
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T22861
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A;Residues: 1-335 <KAW>
A;Residues: 1-335 <KAW>
A;Cross-references: GB:AJ248288; GB:AL096836; NID:g5458960; PIDN:CAB50678.1; PID:g545
A;Experimental source: strain Orsay
C;Genetics:
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C;Superfamily: conserved
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A; Introns:
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rons: 65/2;
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8; Conserv
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100.0%; Pr
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    score 8; DB 2
; Pred. No. 3.7
0; Mismatches
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s; Pred. No. 3.5
0; Mismatches
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                       7.2
    0;
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RESULT 3
E97011
ATPase with
C; Species: C
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T38482
                C; Accession: T38482
R; Brown, D.; Churcher, C.M.; Wood, V.; Barrell, submitted to the EMBL Data Library, June 1997
A; Reference number: Z21796
A; Accession: T38482
                                                                                                          hypothetical protein SPAC29A4.07 - fission yeast (; Species: Schizosaccharomyces pombe C;Date: 03-Dec-1999 #sequence_revision 03-Dec-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R;Moore, C.A.; Kittilson, J.D.; Dahl, S.K.; Sheridan, M.A. Gen. Comp. Endocrinol. 98, 253-261, 1995
A;Title: Isolation and characterization of a cDNA encoding A;Reference number: I51064; MUID:95354921; PMID:7628684
A;Accession: I51064
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R;Nolling, J.; Breton, G.; Omelchenko, M.V.; Markarova, K.; Daly, M.J.; Bennett, G.N.; Koonin, E.V.; Smith, D.R. J. Bacteriol. 183, 4823-4838, 2001

A;Title: Genome Sequence and Comparative Analysis of the A;Reference number: A96900; MUID:21359325; PMID:21359325

A;Accession: E97011
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C:Species: Oncorhynchus mykiss (rainbow trout)
C:Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       somatostatin II precursor - rainbow trout C; Species: Oncorhynchus mykiss (rainbow to
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C;Superfamily: endopeptidase Clp ATP-binding chain
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C; Date: 14-Sep-2001 #sequence_revision 14-Sep-2001 #text_change 30-Sep-2001
C; Accession: E97011
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A; Residues: 1-752 < KUR>
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A; Status: preliminary; translated from GB/EMBL/DDBJ
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Pred. No.
                                                                                                                                                 fission yeast (Schizosaccharomyces pombe)
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A; Molecule type: protein
A; Residues: 17-46 <HOK>
C; Comment: This protein is used to stabilize the displaced single strand of of C; Comment: This protein is used to stabilize the displaced single strand of of C; Comment: This protein is used to stabilize the displaced single stranded DNA-binding protein; single-stranded DNA-binding protein; single-stranded mitochondrion) #status predicted <SIG>
F; 11-148/Product: single-stranded mitochondrial DNA-binding protein #status predicted <SIG>
F; 46-130/Domain: single-stranded DNA-binding protein homology <SSD>
                                                                                                                                                                                                                              R;Gattung, S.; Scheel, r. submitted to the EMBL Data Library, July 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Uross-references: EMBL:Z97210; PIDN:CAB10134.1; A;Experimental source: strain 972h-; cosmid c29A4 C;Genetics:
                                                                                                                                                              A; Description: The sequence of A; Reference number: Z21105 A; Accession: T31967
                                                                                                                                                                                                                                                                                                                                                                              hypothetical protein C33C12.2 - Caenorhabditis C; Species: Caenorhabditis elegans C; Date: 29-Oct-1999 #sequence_revision 29-Oct-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Arch. Biochem. Biophys. 282, 116-124, 1990
A;Title: Structural and functional studies of the rat mitochondrial single A;Reference number: S13264; MUID:91024186; PMID:2221914
A;Accession: S13264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gene 126, 219-225, 1993
A;Title: Cloning of human and rat cDNAs encoding the mitochondrial single-stranded A;Reference number: JN0568; MUID:93246247; PMID:8482537
A;Accession: JN0569
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A;Gene: SPDB:SPAC29A4.07
A;Map position: 1
                                                                   A; Molecule type: DNA
A; Residues: 1-170 <GAT>
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A; Residues: 1-151 <TIR>
A; Cross-references: GB: M94557;
A; Note: the authors translated
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A; Residues: 1-136 <BRO>
A;Cross-references: EMBL:AF016662; PIDN:AAB66062.1; GSPDB:GN00020; CESP:C33C12.2
A;Experimental source: strain Bristol N2; clone C33C12
                                                                                                                             A; Status: preliminary; translated
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R;Chambaud, I.; Heilig, R.; Ferris, S.; Barbe, V.; Samson, D.; Galisson, F.; Moszer, I.; Nucleic Acids Res. 29, 2145-2153, 2001
A;Title: The complete genome sequence of the murine respiratory pathogen Mycoplasma pulm A;Reference number: A99512; MUID:21267165; PMID:11353084
A;Recession: D90569
                                                                                                                                                                                                                                                                                                                                            D: Jones, L.M.; Karst, U.
Science 294, 849-852, 2001
A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; Madueno, E.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-Boland, J.A.; A; Title: Comparative genomics of Listeria species.
A; Reference number: AB1077; MUID:21537279; PMID:11679669
A; Accession: AB1527
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A;Gene: CESP:C33C12.2
A;Map position: 2
A;Introns: 69/3; 113/2
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                                                                                                                                                                                                                                   A;Cross-references: GB:AL592022; PIDN:CAC95986.1; PID:g16413205; GSPDB:GN00178 A;Experimental source: strain Clip11262
                                                                                                                                                                                                                                                                                A; Molecule type: DNA
A; Residues: 1-202 <GLA>
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A:Genetic code: SGC3
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A; Residues: 1-180 <KUR>
A; Cross-references: GB:AL445566; PID:g14089874; PIDN:CAC13633.1; GSPDB:GN00153
A; Experimental source: strain UAB CTIP
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                                                     HDGSLDL 177
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FLIEKLI 62
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Fsihi, H.
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Wehland,
hypothetical protein MJ0042 - Methanococcus jannaschii C;Species: Methanococcus jannaschii C;Species: Methanococcus jannaschii C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 22-Oct-1999 C;Accession: B64305 C;Accession: Mite, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Blab; R;Bult, C.J.; White, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Blab; Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek, rson, J.D.; Sadow, P.W.; Hanna, M.C.; Cotton, M.D.; Roberts, K.M.; Hurst, M.A. Science 273, 1058-1073, 1996 A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.; Woese A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannasc
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C; Date: V. ..... C; Accession: $49599
C; Accession: $49599
R; Immonen, T.; Paulin, L.; Saris, P.
submitted to the EMBL Data Library, January 1994
submitted to the EMBL Data Library and the conjugative nisin-sucrose
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Science 294, 849-852, 2001
A; Authors: Kreft, J.; Kuhn, M.; Kunst, F.; Kurapkat, G.; ok, C.; Schlueter, T.; Simoes, N.; Tierrez, A.; Vazquez-IA; Title: Comparative genomics of Listeria species.
A; Reference number: AB1077; MUID:21537279; PMID:11679669
A; Recession: AH1169
                                                                                                                                    A; Reference number: S49597
A; Accession: S49599
A; Status: preliminary
A; Molecule type: DNA
A; Residues: 1-214 < IMM>
A; Cross-references: EMBL: Z29363; NID: g575300; PIDN: CAA82549.1;
C; Superfamily: Lactococcus lactis probable permease nisG
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                                                                                                                                                                                                                                                                                                                                                                  probable permease nisG - Lactococcus lactis
C; Species: Lactococcus lactis
C; Date: 05-Mar-1995 #sequence_revision 12-May-1995
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R;Glaser, P.; Frangeul, L.;
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A;Molecule type: DNA
A;Residues: 1-202 <GLA>
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100.0%; Pr
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                                                                     Score 7; DB 2; Pred. No. 27; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 7; DB 2; Pred. No. 25; 0; Mismatches
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. 27;
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, A.; Vazquez-Boland, J.A.;
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Dussurget, O.;
                                                                                                                                                                                                                                                                                                                                                                     #text_change 15-Oct-1999
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                                                                                                        Length 214;
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                                                                                                                                                                PID:g575303
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        27-Nov-2001
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Voss, H.; Wehla
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Fsihi,
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Woese

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A;Gene: devA
C;Superfamily
C;Keywords: A
F;26-222/Doma
F;43-50/Regio
                                                                                                                                                                                                                                                                           heterocyst maturation protein devA - An C; Species: Anabaena sp. C; Date: 20-Feb-1995 #sequence_revision C; Accession: A55541; S38423 R; Maldener, I.; Fiedler, G; Ernst, A.; J. Bacteriol. 176, 7543-7549, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT
D64498
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A; Reference number: A64300; MUI
A; Accession: B64305
A; Status: preliminary; nucleic
A; Molecule type: DNA
A; Residues: 1-215 < BUL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C; Species: Methanococcus jannaschii
C; Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 21-Jul-2000
C; Accession: D64498
R; Bult, C.J.; White, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Blal; Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek,
                                                                                                                                                                  A;Cross-references: EMBL:X75422; NID:g407772; PIDN:CAA53174.1; C;Genetics:
                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-244 <MAL>
                                                                                                                                                                                                                                   A;Title: Characterization of devA, a gene required for the maturation of proheterocysts A;Reference number: A55541; MUID:95095923; PMID:8002578 A;Accession: A55541
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A; Residues: 1-241 <BUL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rson, J.D.; Sadow, P.W.; Hanna, M.C.; Cotton, M.D.; Roberts, K.M.; Hurst, M.A. Science 273, 1058-1073, 1996
A; Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ş
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Status: preliminary; nucleic acid sequence not shown; translation
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Best Local S
Matches 7
                                                                            Superfamily: short-chain ATP-binding cassette proteins; Keywords: ATP; nucleotide binding; P-loop; 26-222/Domain: ATP-binding cassette homology <ABC>; 43-50/Region: nucleotide-binding motif A (P-loop)
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                                 Local
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                 Similarity 7; Conserv
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                  Conservative
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            3./
100.0%; F,
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                              Score 7; L
Pred. No.
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Pred. No.
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27;
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29;
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               0;
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                                                Length
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                                                                                                                                  ATP-binding
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               0;
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dek, A.
            translation initiation factor IF-2, chloroplast - r C; Species: chloroplast Cyanidium caldarium C; Date: 06-Jan-1995 #sequence_revision 24-Oct-1997 C; Accession: S36410; S70822 R; Kostrzewa, M.; Zetsche, K.
submitted to
                                                                                                    S36410
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Best Local
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A;Note: Nostoc sp. scram. C;Date: 14-Dec-2001 #sequence_revision 14 C;Accession: AI2269 R;Kaneko, T.; Nakamura, Y.; Wolk, C.P.; K
                                                                                                                                                                                                                                                     R;Chambaud, I.; Heilig, R.; Ferris, S.; Barbe, V.; Samson, D.; Nucleic Acids Res. 29, 2145-2153, 2001
A;Title: The complete genome sequence of the murine respiratory A;Reference number: A99512; MUID:21267165; PMID:11353084
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Gene: devA
C;Superfamily: short-chain ATP-binding cassette proteins; ATP-binding
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DNA Res. 8, 205-213, 2001
A; Title: Complete Genomic
                                                                                                     A;Gene: MYPU_1210
A;Genetic code: SGC3
                                                                                                                                                                                      A; Molecule type: DNA
A; Residues: 1-245 < KUR>
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C:Accession: A90527
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A; Residues: 1-244 < KUR>
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                                                                                                                                                      A;Cross-references: GB:AL445566; PID:g14089534; PIDN:CAC13294.1; A;Experimental source: strain UAB CTIP
                                                                                                                                                                                                                      A; Status: preliminary
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                                                                                                                                                                                                                                                                                                                                                                            hypothetical protein MYPU_1210 [imported] - Mycoplasma pulmonis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   181 RSVQEGN 187
96 TLIVLII 102
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                                   Similarity 7; Conser
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001 #sequence_revision 14-Dec-2001
                               3.7%;
ilarity 100.0%;
Conservative
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                                  Score 7; DB 2; Pred. No. 30; 0; Mismatches
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Pred. No.
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; Takazawa, 1
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                                         DB ,
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; Sasamoto,
M.; Yamada,
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#text_change
                                                                 Length 245;
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                                   Indels
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30-Jun-2002
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Data Library,

June 1992

red alga (Cyanidium caldarium)

#text_change

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UDP-glucose dehydrogenase (ugd-1) homolog - Archaeoglobus fulgidus
C; Species: Archaeoglobus fulgidus
C; Species: Archaeoglobus fulgidus
C; Date: 05-Dec-1997 #sequence_revision 05-Dec-1997 #text_change 11-Jun-1999
C; Accession: F69287
R; Klenk, H.P.; Clayton, R.A.; Tomb, J.F.; White, O.; Nelson, K.E.; Ketchum, K.A.; Dodson
.; Fleischmann, R.D.; Quackenbush, J.; Lee, N.H.; Sutton, G.G.; Gill, S.; Kirkness, E.F.
Glodek, A.; Zhou, L.; Overbeek, R.; Gocayne, J.D.; Weidman, J.F.; McDonald, L.
Nature 390, 364-370, 1997
A; Authors: Utterback, T.; Cotton, M.D.; Spriggs, T.; Artiach, P.; Kaine, B.P.; Sykes, S.
Smith, H.O.; Woese, C.R.; Venter, J.C.
A; Title: The complete genome sequence of the hyperthermophilic, sulfate-reducing archaec
A; Reference number: A69250; MUID:98049343; PMID:9389475
A; Accession: F69287
A; Status: preliminary; nucleic acid sequence not shown; translation not shown
A; Molecule type: DNA
A; Residues: 1-324 < KLE>
A; Cross-references: GB:AE001084; GB:AE000782; NID:g2689407; PIDN:AAB90929.1; PID:g265033
C; Superfamily: GDPmannose dehydrogenase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Reference number: S36410
A; Accession: S36410
A; Molecule type: DNA
A; Residues: 1-259 < KOS>
A; Cross-references: EMBL: X66698; NID: g396522; PID: g396523
A; Cross-reference: Strain 14-1-1
R; Kostrzewa, M.; Zetsche, K.
Plant Mol. Biol. 23, 67-76, 1993
hypothetical protein Y94A7B.5 - Caenorhabditis elegans C; Species: Caenorhabditis elegans C; Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #te: C; Accession: T27484 R; Kershaw, J. submitted to the EMBL Data Library, June 1998 A; Reference number: Z20373 A; Accession: T27484
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Genome: chloroplast
C; Superfamily: translatio
C; Keywords: chloroplast;
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A;Experimental source: strain 14-1-1
A;Note: the source is designated as Galdieria sulphuraria
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37;
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A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-327 <WIL>
A; Cross-references: EMBL:AL023856; PIDN:CAA19567.1; GS
A; Experimental source: clone Y94A7B
C; Genetics:
A; Gene: CESP:Y94A7B.5
A; Map position: 5
A; Introns: 104/1; 245/3
C; Superfamily: Caenorhabditis hypothetical protein C45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT
S56333
                   A;Cross-references: EMBL:U14003; NID:g1263172; PIDN:AAA97004.1; PID:g536949
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, Augu R;Blattner, F.R.; Plunkett III, G.; Bloch, C.A.; Perna, N.T.; Burland, V.; F.A.; Rose, D.J.; Mau, B.; Shao, Y.
Science 277, 1453-1462, 1997
A;Title: The complete genome sequence of Escherichia coli K-12.
A;Reference number: A64720; MUID:97426617; PMID:9278503
                                                                                                                                                                                                                                                                                              carbon-phosphorus lyase phnD - Escherichia coli (strain K-12)
C;Species: Escherichia coli
C;Date: 28-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 01-Mar-
C;Date: 28-Oct-1995 #sequence_revision 03-Nov-1995 #text_change 01-Mar-
C;Accession: S56333, H65219; E35718
R;Burland, V.; Plunkett III, G.; Sofia, H.J.; Daniels, D.L.; Blattner,
Nucleic Acids Res. 23, 2105-2119, 1995
A;Title: Analysis of the Escherichia coli genome VI: DNA sequence of the A;Reference number: S56314; MUID:95334362; PMID:7610040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    conserved hypothetical protein BB0619 - Lyme disease spirochete C; Species: Borrelia burgdorferi (Lyme disease spirochete) C; Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 21-Jul-2000 C; Date: 15-Oct-1999 #sequence_revision 15-Oct-1999 #text_change 21-Jul-2000 C; Date: 190177 C; Date: 190177 C; Date: 190177 C; Date: 190177 C; Casjens, S.; Huang, W.M.; Sutton, G.G.; Clayton, R.; Lathigra, Son, D.; Peterson, J.; Kerlavage, A.R.; Quackenbush, J.; Salzberg, S.; Hanson, Bowman, C.; Garland, S.; Fujii, C.; Cotton, M.D.; Horst, K.; Roberts, K.; Hature 390, 580-586, 1997
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B70177
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary; nucleic acid sequence not shown; translation A;Molecule type: DNA A;Residues: 1-330 <KLE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Nature 390, 580-586, 1997
A; Authors: Smith, H.O.; Venter, J.C.
A; Title: Genomic sequence of a Lyme disease spirochaete, A; Reference number: A70100; MUID:98065943; PMID:9403685
A; Accession: B70177
                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-338 <BUR>
A; Reference number: A; Accession: H65219
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100.0%; Pr
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                                                                                                                          Riley, M.;
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son, M.;
Hatch,
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B 전화

acid

sequence

not

shown;

translation not shown

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A; Molecule type: DNA
A; Residues: 1-338 <BLAT>
A; Residues: 1-338 <BLAT>
A; Cross-references: GB: AE000482; GB: U00096; NID: g2367349; PIDN: AAC77066.1;
A; Experimental source: strain K-12, substrain MG1655
R; Chen, C.M.; Ye, Q.Z.; Zhu, Z.; Wanner, B.L.; Walsh, C.T.
J. Biol. Chem. 265, 4461-4471, 1990
A; Title: Molecular biology of carbon-phosphorus bond cleavage. Cloning and A; Reference number: A35718; MUID: 90170953; PMID: 2155230
                                                       A;Cross-references: GB:AE005174; NID:g12519072; PIDN:AAG59304.1; GSPDB:GN00145; UWGP:Z5:A;Experimental source: strain O157:H7, substrain EDL933 C;Genetics:
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D86105
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A; Molecule 1-338 <HAY>
A; Residues: 1-338 <HAY>
A; Cross-references: GB:BA000007; PIDN:BAB38510.1; PID:g13364564;
A; Cross-reference: strain O157:H7, substrain RIMD 0509952
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                                                                                                                                                                                                                                                         R;Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasiiller, L.; Grotbeck, E.J.; Davis, N.W.; Lim, A.; Dimalanta, Nature 409, 529-533, 2001
A;Title: Genome sequence of enterohemorrhagic Escherichia coa;Reference number: A85480; MUID:21074935; PMID:11206551
                                                                                                                                                                                                                                                                                                                                                                     hypothetical protein phnD [imported] - Escherichia coli (strain O157:H7, su C; Species: Escherichia coli (c; Date: 16-Feb-2001 #sequence_revision 16-Feb-2001 #text_change 14-Sep-2001 C; Accession: D86105 R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Burland, V.; Mau, B.; Glasner, J.D.; Rose, R; Perna, N.T.; Plunkett III, G.; Perna, R; Perna, R;
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A;Title: Complete genome sequence of enterohemorrhagic Escherichia coli O157:H7
A;Reference number: A99629; MUID:21156231; PMID:11258796
                                                                                                                                            A; Molecule type: DNA
A; Residues: 1-338 <STO>
                                                                                                                                                                                                                             A; Reference number: A85480; A; Accession: D86105
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A; Residues: 1-311,'A',313-338
A; Cross-references: GB:J05260;
                                                                                                                                                                                                 A; Status: preliminary
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Kasunaga, T.; Kuhara,
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100,08; 27
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'; NID:g147192; PIDN:AAA24340.1; PID:g147197
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Shiba, T.;
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; Hattori, I
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lanta, E.;
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M.; Shinagawa,
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K.; A
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Apodaca,
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A; Note: ND5
C; Superfamil
C; Keywords:
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Gene 203,
                                                                                                                                                                                                                                                                                                                                                                                                         NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 (isolate High Tatras Slovakia) C;Species: mitochondrion Carabus linnei C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 03-Jun-2002 C;Accession: T12104; T12106 R;Imura, Y.; Su, Z.H.; Kim, C.G.; Osawa, S. Elytra 26, 223-248, 1998
                                                                                                                                                                                                                                                                                                                                                           A;Title: Reorganization of the Oreocarabus complex (Coleoptera, A;Reference number: Z17381 A;Accession: T12104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: EMBL:AF007101;
A;Experimental source: ATCC 29253
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    probable pteridine-dependent dioxygenase - Streptomyces hygroscopicus
C;Species: Streptomyces hygroscopicus
C;Date: 24-Mar-1999 #sequence_revision 24-Mar-1999 #text_change 22-Oct-1999
                                                                                                                                                                                 A; Experimental source: isolate Zilina
                                                                                                                                                                                             A; Molecule type: DNA
A; Residues: 1-122, 'L', 124-355 <IM2>
A; Cross-references: EMBL: AB017473;
                                                                                                                                                                                                                                                                           A;Cross-references: EMBL:AB017472; PIDN:BAA33188.1
A;Experimental source: isolate High tatras Slovakia;
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A; Residues: 1-355 < IMU>
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                                                                                                                                                                                                                                            A; Status: preliminary; translated
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Best Local
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Best Local :
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Best Local (
                                                                                                               Superfamily: NADH dehydrogenase (ubiquinone) chain
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20 FIFLLFF 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         X.; Stassi, D.; Lax, S.; Katz, 3, 1-9, 1997
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                                                                                              complex;
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                                                                                                                                                                               PIDN:BAA33189.1 ina Slovakia; adult;
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R;Su, Z.H.; Tominaga, O.; Okamoto, M.; Osawa, S.
Mol. Biol. Evol. 15, 1026-1039, 1998
A;Title: Origin and diversification of hindwingless Damaster ground beetles within A;Reference number: Z17319; MUID:98384842; PMID:9718730
A;Status: preliminary: +----
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C; Superfamil
C; Keywords:
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R;Su, Z.H.; Tominaga, O.; Okamoto, M.; Osawa, S.
Mol. Biol. Evol. 15, 1026-1039, 1998
A;Title: Origin and diversification of hindwingless Damaster A;Reference number: Z17319; MUID:98384842; PMID:9718730
A;Accession: T13831
A;Status: preliminary; translated from GR/FMRT./DDBT
                                                                                       A;Cross-references: EMBL:D50339; NID:g1000905; PIDN:BAA08874.1; A;Experimental source: adult; thorax muscle C;Genetics:
A;Genome: mitochondrion
A;Genome: mitochondrion
A;Note: ND5
C;Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C;Keywords: membrane-associated complex; mitochondrion; NAD; oxi
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A; Experimental sourc
C; Genetics:
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C; Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 03-Jun-2002
C; Accession: T13830
R; Su, Z.H.; Tominaga, O.; Okamoto, M.; Osawa, S.
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A; Residues: 1-355 <SUZ>
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cies: mitochondrion Acoptolabrus gehinii
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membrane-associated complex; mitochondrion; NAD; oxidative
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Khumaratkhok Russia; adult; thorax muscle
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A; Reference number: Z17787; MUID:97077297; PMID:891
A; Accession: T13833
A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: DNA
A; Residues: 1-355 <SUZ>
                                                                                                                                                                                                     R;Su, Z.H.; Ohama, T.; Okada, T.S.; Nakamura, K.; Ishikawa, R.; Osawa, S. J. Mol. Evol. 42, 124-129, 1996
A;Title: Phylogenetic relationships and evolution of the Japanese Carabinae A;Reference number: Z17787; MUID:97077297; PMID:8919864
                                                                                                                                                                                                                                                                                     NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - Asthenocarabus opace C; Species: mitochondrion Asthenocarabus opaculus C; Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 03-Jun-2002 C; Accession: T13833
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Note: ND5
C; Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C; Keywords: membrane-associated complex; mitochondrion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-355 <IMU>
A; Cross-references: EMB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            R; Imura, Y.; Zhou, H.Z.; Okamoto, M.; Su, Z.H.; Osawa, S. submitted to the EMBL Data Library, January 1998 a; Description: Phylogenetic relationships of some Chinese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - Aristocarabus viridifossulatu C;Species: mitochondrion Aristocarabus viridifossulatus C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 03-Jun-2002 C;Accession: T11601
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C;Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C;Keywords: membrane-associated complex; mitochondrion; NAD; oxidative phosphorylatio
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A; Residues: 1-355 <SUZ>
A; Cross-references: EMBL:D50429; PIDN:BAA08925.1
A; Experimental source: adult; thorax muscle
C; Genetics:
     C; Superfami
C; Keywords:
                                                            A; Experimental source: C; Genetics:
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C;Genetics:
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A; Status: preliminary; translated
                                                                                                      A; Cross-references:
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                                          A; Genome: mitochondrion
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Superfamily: NADH u
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                                                                                   EMBL:D50340; NID:g1000901; PIDN:BAA08875.1; ce: adult; thorax muscle
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ate Baoxing-Sichuan;
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                                                                                                      PID:g1438079
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Similarity

3.7%; 100.0%;

Score 7; Pred. No.

DB 2;

Length

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T12029

NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - C; Species: mitochondrion Carabus concolor C; Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text C; Accession: T12029
                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Su, Z.H.; Ohama, T.; Okada, T.S.; Nakamura, K.; Ishikawa, R.; Osawa, S. J. Mol. Evol. 42, 124-129, 1996
A;Title: Phylogenetic relationships and evolution of the Japanese Carabinae A;Reference number: Z17787; MUID:97077297; PMID:8919864
A;Accession: T13862
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A;Cross-references: EMBL:D50343; NID:g1000915; PIDN:BAA08878.1; A;Experimental source: adult; thorax muscle
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C;Accession: T13861
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A;Title: Phylogenetic relationships and evolution of the Japanese Carabinae A;Reference number: Z17787; MUID:97077297; PMID:8919864
A;Accession: T13861
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A; Cross-references: EMBJ
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A; Residues: 1-355 <S
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                                                                                                                                                                                                                                                                             Superfamily: NADH dehydrogenase (ubiquinone) chain 5; Keywords: membrane-associated complex; mitochondrion;
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cies: mitochondrion Campalita chinense
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                                                                                                                                                                                                                                           Length 355;
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 A; Experimental source: C; Genetics:
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Elytra 26, 223-248, 1998
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                                                                                                                                                        R; Imura, Y.; Su, Z.H.; K
Elytra 26, 223-248, 1998
                                                                                                                                                                                           C; Accession: T12028
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A; Residues: 1-355 <I
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A; Note: ND5
C; Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C; Keywords: membrane-associated complex; mitochondrion;
                                                                             A;Title: Reorganization of A;Reference number: Z17381 A;Accession: T12028
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A; Accession: T12029
A; Statue
            A; Molecule type: DNA
A; Residues: 1-355 < IMU>
A; Cross-references: EMBL: AB017455; PIDN: BAA33171.1
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A;Experimental source: isolate Artvin Turkey; adult;
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                                                            A; Status: preliminary; translated
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ytra 26, 223-248, 1998
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Champorcer Italy No1;
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NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - Cara C;Species: mitochondrion Carabus hortensis C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_cha C;Accession: T12034
R;Imura, Y; Su, Z.H.; Kim, C.G.; Osawa, S.
Elytra 26, 223-248, 1998
A;Title: Reorganization of the Oreocarabus complex (Coleopte A;Reference number: Z17381
A;Reference number: Z17381
A;Resion: T12034
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-355 <IMU>
A;Cross-references: EMBL:AB017463; PIDN:BAA33179.1
A;Experimental source: isolate Innsbruck Austria; adult; tho C;Genetics:
A;Genome: mitochondrion
A;Note: ND5
C;Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C;Keywords: membrane-associated complex; mitochondrion; NAD;
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C; Superfami
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A;Note: ND5
C;Superfamily: NADH deh
C;Keywords: membrane-as:
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A;Cross-references: EMBL:AB017450;
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A; Residues: 1-355 < IMU>
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ilarity 100.0%;
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d; Pred. No. 40;
0; Mismatches
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A; Note: ND5
C; Superfamily: |
C; Keywords: meml
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T12114
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T12111
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C; Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C; Keywords: membrane-associated complex; mitochondrion;
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R; Imura, Y.; Su,
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A;Molecule type: DNA
A;Residues: 1-355 <IMU>
A;Cross-references: EMBL:AB017476;
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A; Residues: 1-355 < IMU>
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NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - Carabus C; Species: mitochondrion Carabus porrecticollis kansaiensis C; Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change C; Accession: T12119
R; Kim, C.G.; Su, Z.H.; Osawa, S.
submitted to the EMBL Data Library, January 1999
A; Description: Origin and diversification of Euleptocarabus porr A; Reference number: Z17426
A; Accession: T12119
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C; Superfamil
C; Keywords:
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R;Kim, C.G.; Tominaga, O.; Su, Z.H.; Osawa, S. submitted to the EMBL Data Library, January 1999 submitted to the EMBL Data Library, January 1999
A;Description: Origin and diversification of Euleptocarabus porrecticollis (Coleoptera,
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T12119
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R;Imura, Y.; Su, Z.H.; Kim, C.G.; Osawa, S.
Elytra 26, 223-248, 1998
A;Title: Reorganization of the Oreocarabus complex (Coleoptera, A;Reference number: Z17381
A;Accession: T12033
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A; Accession: T12118
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A; Residues: 1-355 <II
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A; Residues: 1-355 <K
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                  Status: preliminary;
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A; Note: ND5
C; Superfamil
C; Keywords:
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C; Superfamily: NADH dehydrogenase (ubiquinone) chain
C; Keywords: membrane-associated complex; mitochondric
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A;Cross-references: EMBL:AB022571; PIDN:CAB19797.1
A;Experimental source: collection site: kaga Ishikawa
                                                                                                          C; Genetics:
                                                                                                                         A;Cross-references: EMBL:AB017453; A;Experimental source: isolate Mt.
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A; Residues: 1-355 < IMU>
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NADH2 dehydrogenase (ubiquinone) (EC 1.6.5.3) chain 5 - Carabus putzeysianus mitochondri C;Species: mitochondrion Carabus putzeysianus raynaudianus C;Date: 05-Nov-1999 #sequence_revision 05-Nov-1999 #text_change 03-Jun-2002 C;Accession: T12154
R;Imura, Y; Su, Z.H.; Kim, C.G.; Osawa, S.
Elytra 26, 223-248, 1998
A;Title: Reorganization of the Oreocarabus complex (Coleoptera, Carabidae) based on endc A;Reference number: Z17381
A;Accession: T12154
A;Status: preliminary; translated from GB/EMBL/DDBJ
A;Cross-references: EMBL:AB017448; PIDN:BAA33164.1
A;Experimental source: isolate Laghi di Lollalunga Italy; adult; thorax muscle C;Genetics:
A;Genome: mitochondrion
A;Note: ND5
C;Superfamily: NADH dehydrogenase (ubiquinone) chain 5
C;Keywords: membrane-associated complex; mitochondrion; NAD; oxidative phosphorylation;
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